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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/924,125

DATE: 12/13/2001
TIME: 13:58:26

Input Set : A:\ES.txt
Output Set: N:\CRF3\12132001\I924125.raw

3 <110> APPLICANT: Communi, Didier
5 <120> TITLE OF INVENTION: THE NATURAL LIGAND FOR ORPHAN G PROTEIN COUPLED RECEPTOR
GPR86 AND

6 METHODS OF USE

8 <130> FILE REFERENCE: 9049/2092

10 <140> CURRENT APPLICATION NUMBER: 09/924,125

11 <141> CURRENT FILING DATE: 2001-07-08

13 <150> PRIOR APPLICATION NUMBER: US 09/924,125

14 <151> PRIOR FILING DATE: 2001-07-08

16 <160> NUMBER OF SEQ ID NOS: 9

18 <170> SOFTWARE: PatentIn version 3.1

20 <210> SEQ ID NO: 1

21 <211> LENGTH: 1002

22 <212> TYPE: DNA

23 <213> ORGANISM: Homo sapiens

25 <400> SEQUENCE: 1

26 atgaacacca cagtgatgca aggcttcaac agatctgagc ggtgccccag agacactcgg 60
28 atagtacagc tggattccc agccctctac acagtgggtt tcttgaccgg catcctgctg 120
30 aatactttgg ctctgtgggt gttgtttcac atccccagct cctccacctt catcatctac 180
32 ctcaaaaaca ctttggtggc cgacttgata atgacactca tgcttccttt caaaatcctc 240
34 tctgactcac acctggcacc ctggcagctc agagcttttg tgtgtcgttt ttcttcggtg 300
36 atattttatg agaccatgta tgtgggcacg gtgctgtag ggctcatagc ctttgacaga 360
38 ttctcaaga tcatcagacc tttagaaaat atttttctaa aaaaacctgt ttttgcaaaa 420
40 acggtctcaa tcttcatctg gttctttttg ttcttcatct ccctgccaaa tatgatcttg 480
42 agcaacaagg aagcaacacc atcgtctgtg aaaaagtgtg ctctcttaaa ggggcctctg 540
44 gggctgaaat ggcacaaat ggtaaataac atatgccagt ttattttctg gactgttttt 600
46 atcctaatagc ttgtgtttta tgtggttatt gcaaaaaaag tatatgattc ttatagaaag 660
48 tccaaaagta aggacagaaa aaacaacaaa aagctggaag gcaaagtatt tgtgtcgtg 720
50 gctgtcttct ttgtgtgttt tgcctcattt cattttgccg gagttccata tactcacagt 780
52 caaaccaaca ataagactga ctgtagactg caaaatcaac tgtttattgc taaagaaaca 840
54 actctctttt tggcagcaac taacatttgt atggatccct taatatacat attcttatgt 900
56 aaaaaattca cagaaaagct accatgtatg caagggagaa agaccacagc atcaagccaa 960
58 gaaaatcata gcagtcagac agacaacata accttaggct ga 1002

61 <210> SEQ ID NO: 2

62 <211> LENGTH: 333

63 <212> TYPE: PRT

64 <213> ORGANISM: Homo sapiens

66 <400> SEQUENCE: 2

68 Met Asn Thr Thr Val Met Gln Gly Phe Asn Arg Ser Glu Arg Cys Pro
69 1 5 10 15
72 Arg Asp Thr Arg Ile Val Gln Leu Val Phe Pro Ala Leu Tyr Thr Val
73 20 25 30
76 Val Phe Leu Thr Gly Ile Leu Leu Asn Thr Leu Ala Leu Trp Val Phe
77 35 40 45
80 Val His Ile Pro Ser Ser Ser Thr Phe Ile Ile Tyr Leu Lys Asn Thr
81 50 55 60
84 Leu Val Ala Asp Leu Ile Met Thr Leu Met Leu Pro Phe Lys Ile Leu
85 65 70 75 80

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88 Ser Asp Ser His Leu Ala Pro Trp Gln Leu Arg Ala Phe Val Cys Arg
89           85           90           95
92 Phe Ser Ser Val Ile Phe Tyr Glu Thr Met Tyr Val Gly Ile Val Leu
93           100          105          110
96 Leu Gly Leu Ile Ala Phe Asp Arg Phe Leu Lys Ile Ile Arg Pro Leu
97           115          120          125
100 Arg Asn Ile Phe Leu Lys Lys Pro Val Phe Ala Lys Thr Val Ser Ile
101           130          135          140
104 Phe Ile Trp Phe Phe Leu Phe Phe Ile Ser Leu Pro Asn Met Ile Leu
105 145           150          155          160
108 Ser Asn Lys Glu Ala Thr Pro Ser Ser Val Lys Lys Cys Ala Ser Leu
109           165          170          175
112 Lys Gly Pro Leu Gly Leu Lys Trp His Gln Met Val Asn Asn Ile Cys
113           180          185          190
116 Gln Phe Ile Phe Trp Thr Val Phe Ile Leu Met Leu Val Phe Tyr Val
117           195          200          205
120 Val Ile Ala Lys Lys Val Tyr Asp Ser Tyr Arg Lys Ser Lys Ser Lys
121           210          215          220
124 Asp Arg Lys Asn Asn Lys Lys Leu Glu Gly Lys Val Phe Val Val Val
125 225           230          235          240
128 Ala Val Phe Phe Val Cys Phe Ala Pro Phe His Phe Ala Arg Val Pro
129           245          250          255
132 Tyr Thr His Ser Gln Thr Asn Asn Lys Thr Asp Cys Arg Leu Gln Asn
133           260          265          270
136 Gln Leu Phe Ile Ala Lys Glu Thr Thr Leu Phe Leu Ala Ala Thr Asn
137           275          280          285
140 Ile Cys Met Asp Pro Leu Ile Tyr Ile Phe Leu Cys Lys Lys Phe Thr
141           290          295          300
144 Glu Lys Leu Pro Cys Met Gln Gly Arg Lys Thr Thr Ala Ser Ser Gln
145 305           310          315          320
148 Glu Asn His Ser Ser Gln Thr Asp Asn Ile Thr Leu Gly
149           325          330

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152 <210> SEQ ID NO: 3

153 <211> LENGTH: 11

154 <212> TYPE: DNA

C--> 155 <213> ORGANISM: Artificial

157 <220> FEATURE:

158 <221> NAME/KEY: misc_binding

159 <222> LOCATION: (1)..(11)

160 <223> OTHER INFORMATION: NF-kB binding element

163 <400> SEQUENCE: 3

164 ggggactttc c

11

167 <210> SEQ ID NO: 4

168 <211> LENGTH: 31

169 <212> TYPE: DNA

C--> 170 <213> ORGANISM: Artificial

172 <220> FEATURE:

W--> 173 <221> NAME/KEY: primer

174 <222> LOCATION: (1)..(31)

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175 <223> OTHER INFORMATION: GPR86 human receptor: a sense primer
178 <400> SEQUENCE: 4
179 ccggaattca ccataaacac cacagtgatg c 31
182 <210> SEQ ID NO: 5
183 <211> LENGTH: 31
184 <212> TYPE: DNA
C--> 185 <213> ORGANISM: Artificial
187 <220> FEATURE:
W--> 188 <221> NAME/KEY: primer
189 <222> LOCATION: (1)..(31)
190 <223> OTHER INFORMATION: GPR86 human receptor: anti-sense primer
193 <400> SEQUENCE: 5
194 cttgtctaga tcagcctaag gttatgttgt c 31
197 <210> SEQ ID NO: 6
198 <211> LENGTH: 20
199 <212> TYPE: DNA
C--> 200 <213> ORGANISM: Artificial
202 <220> FEATURE:
W--> 203 <221> NAME/KEY: primer
204 <222> LOCATION: (1)..(20)
205 <223> OTHER INFORMATION: GPR86 sense primer
208 <400> SEQUENCE: 6
209 tgtgtcggtt ttcttcggtg 20
212 <210> SEQ ID NO: 7
213 <211> LENGTH: 18
214 <212> TYPE: DNA
C--> 215 <213> ORGANISM: Artificial
217 <220> FEATURE:
W--> 218 <221> NAME/KEY: primer
219 <222> LOCATION: (1)..(18)
220 <223> OTHER INFORMATION: GPR86 antisense primer
223 <400> SEQUENCE: 7
224 ctgccaaaaa gagagttg 18
227 <210> SEQ ID NO: 8
228 <211> LENGTH: 20
229 <212> TYPE: DNA
C--> 230 <213> ORGANISM: Artificial
232 <220> FEATURE:
W--> 233 <221> NAME/KEY: primer
234 <222> LOCATION: (1)..(20)
235 <223> OTHER INFORMATION: aldolase sense primer
238 <400> SEQUENCE: 8
239 ggcaagggca tcctggctgc 20
242 <210> SEQ ID NO: 9
243 <211> LENGTH: 23
244 <212> TYPE: DNA
C--> 245 <213> ORGANISM: Artificial
247 <220> FEATURE:
W--> 248 <221> NAME/KEY: primer

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249 <222> LOCATION: (1)..(23)

250 <223> OTHER INFORMATION: aldolase antisense reverse primer

253 <400> SEQUENCE: 9

254 taacggggcca gaacattggc att

23

VERIFICATION SUMMARY

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Input Set : A:\ES.txt

Output Set: N:\CRF3\12132001\I924125.raw

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L:170 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:173 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:185 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:188 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:200 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:203 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:215 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:218 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7
L:230 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:233 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:245 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:248 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9